

CofS

PATENT Customer No. 22,852 Attorney Docket No. 6832.0013

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re U.S. Patent No.: 6,926,898	1									
Inventors:	1									
Craig A. Rosen and William A. Haseltine										
Issue Date.: August 9, 2005										
For: ALBUMIN FUSION PROTEINS)									
Commissioner for Patents P.O. Box 1450 Alexandria, VA 22313-1450										
Sir:										

REQUEST FOR CERTIFICATE OF CORRECTION

Pursuant to 35 U.S.C. §§ 254 and 255, and 37 C.F.R. §§ 1.322 and 1.323, this is a request for a Certificate of Correction in the above-identified patent. Some of the mistakes identified in the appended Form occurred through the fault of the Patent Office, as clearly disclosed by the records of the application which matured into this patent.

For example, the priority claims to Provisional Application Nos. 60/256,931, filed December 21, 2000; 60/199,384, filed April 25, 2000; and 60/229,358, filed April 12, 2000, were deleted in an Amendment filed February 4, 2004, and a Corrected Filing Receipt reflecting the change was mailed by the PTO on February 13, 2004. However, 189,99 0P the issued patent was printed with the priority claims in the title page under item (60).

Furthermore, the omitted U.S. Patent Documents under item (56) (References Cited) in the title page, were cited by Applicants in an Information Disclosure Statement

filed April 5, 2004, and the Office returned the initialed Form PTO 1449 with the Supplemental Notice of Allowance mailed June 29, 2004.

The omitted OTHER PUBLICATIONS under item (56) (References Cited) in the title page, were also cited by Applicants in the Information Disclosure Statement filed April 5, 2004, and the Office returned the initialed Form PTO 1449 by facsimile on July 23, 2004.

The issued patent was printed without the Examiner's Amendment to the specification mailed March 3, 2005, with the Supplemental Notice of Allowance. The attached Certificate of Correction amends the specification according to the Examiner's Amendment.

Furthermore, the issued patent reflects the original Sequence Listing filed rather than the Substitute Sequence Listing submitted on August 20, 2004. The Sequence Listing in the attached Certificate of Correction is identical to the Substitute Sequence Listing filed on August 20, 2004, and is also identical to the computer readable copy of the Substitute Sequence Listing also submitted on August 20, 2004. Thus, the correction contains no new matter.

Other mistakes identified in the appended Form are of a clerical or typographical nature, or of minor character, and resulted from an error made in good faith by patentees. A check in the amount of \$100 (the fee set forth in 37 C.F.R. § 1.20(a)) is attached. Should a check not be appended or should any additional fees be needed, authorization is hereby given to charge any fees due in connection with the filing of this request to Deposit Account No. 06-0916.

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Two (2) copies of PTO Form 1050 are appended. The complete Certificate of Correction involves thirty-five (35) pages. Issuance of the Certificate of Correction containing the correction is earnestly requested.

Please charge any required fees not included herewith to our deposit account 06-0916.

Respectfully submitted,

FINNEGAN, HENDERSON, FARABOW, GARRETT & DUNNER, L.L.P.

Charle E Van Horn

Dated: December 23, 2005

Charles E. Van Horn Reg. No. 40,266

UNITED STATES PATENT AND TRADEMARK OFFICE CERTIFICATE OF CORRECTION

PATENT NO.

6,926,898

Page 1 of 35

APPLICATION NO.:

09/832,929

ISSUE DATE:

August 9, 2005

INVENTOR(S):

Craig A. Rosen and William A. Haseltine

It is hereby certified that an error or errors appear in the above-identified patent and that said Letters Patent is hereby corrected as shown below:

Under item (60) (Related U.S. Application Data) of the title page, delete the text beginning with "Provisional application No. 60/256,931" to and ending "provisional application No. 60/229,358, filed on Apr. 12, 2000."

Under item (56) (References Cited) of the title page and under U.S. PATENT DOCUMENTS beginning on page 1, insert:

 2003-0022308 A1	1/2003	Fleer et al.
2003-0036170 A1	2/2003	Fleer et al.
2003-0036171 A1	2/2003	Fleer et al.
2003-0036172 A1	2/2003	Fleer et al.
2003-0054554 A1	3/2003	Becquart et al.
2003-0082747 A1	5/2003	Fleer et al.
2003-0104578 A1	10/2001	Ballance
2004-0010134 A1	4/2001	Rosen et al.
09/832,501	4/2001	Ballance et al.
09/833,041	4/2001	Rosen et al.
09/833,111	4/2001	Rosen et al.
09/833,117	4/2001	Rosen et al.
09/833,118	4/2001	Rosen et al.
10/702,536	11/2003	Fleer et al.
10/702,636	11/2003	Fleer et al

MAILING ADDRESS OF SENDER

U.S. Patent No. 6,926,898

Under item (56) (References Cited) of the title page and under OTHER PUBLICATIONS beginning on page 1, insert:

-- Larsson, M., et al., "Role of Annexins in Endocytosis of Antigens in Immature Human Dendritic Cells," *Immunology* 92:501-511 (1997).

Latta, M. et al., "Synthesis and Purification of Mature Human Serum Albumin From E. Coli," Bio/Technology 5:1309-1314 (1987).

Latta, M., et al., "Tryptophan Promoter Derivatives on Multicopy Plasmids: A Comparative Analysis of Expression Potentials in *Escherichia coli*," *DNA and Cell Biology* 9:129-137 (1990).

Lawn, R.M., et al., "The Sequence of Human Serum Albumin cDNA and its Expression in E. coli," *Nucleic Acids Research* 9:6103-6113 (1981).

Le Bras, M., et al., "Epidemiologie et Clinique des Maladies Tropicales D'importation," La Revue de Medicine Interne 13:205-210 (1992), with English translation.

Leblois, H., et al., "Stable Transduction of Actively Dividing Cells via a Novel Adenoviral/Episomal Vector," *Molecular Therapy* 1:314-322 (2000).

Lee, C-H., et al., "Sodium Pertechnetate Tc99m Antral Scan in the Diagnosis of Retained Gastric Antrum," *Arch. Surg.* 119: 309-311 (1984).

Lee, C-L., et al., "Preparation and Characterization of Polyethylene-Glycol-Modified Salmon Calcitonins," *Pharmaceutical Development and Technology*, 4(2): 269-275 (1999).

Lee, W-C., et al., "Identification and Characterization of a Nuclear Localization Sequence-Binding Protein in Yeast," *Proc. Natl. Acad. Sci. USA* 86:8808-8812 (1989).

Lee, Y-H., et al., "Comparison of Effective Renal Plasma Flow (ERPF) and Endogenous Creatinine Clearance (Ccr) in Evaluation of the Differential Kidney Function: An in Vivo Study," *Chin. Med. J. (Taipei)* 49:147-152 (1992).

Lei, H-Y., et al., "An Antigen-specific Hypersensitivity Which Does Not Fit Into Traditional Classification of Hypersensitivity," *The Journal of Immunology* 143:432-438 (1989).

Levitt, D., et al., "Toxicity of Perfluorinated Fatty-Acids for Human and Murine B Cell Lines," *Toxicology and Applied Pharmacology* 86:1-11 (1986).

MAILING ADDRESS OF SENDER

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Lew D.B., et al., "Mitogenic Effect of Lysosomal Hydrolases on Bovine Tracheal Myocytes in Culture," The Journal of Clinical Investigation 88:1969-1975 (1991).
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Lewis, C., et al., "Is Sexual Dysfunctoin in Hypertensive Women Uncommon or Understudied?" *American Jour of Hypertension*," 11:733-735 (1998). --

Under item (57) (ABSTRACT) of the title page, "disordrs" should read --disorders--.

In the Specification

Col. 143, line 26, delete "As exhibited in Table 2, most", and insert -- Most--.

Col. 143, line 31, delete "Table 2".

In the Claims

Col. 340, line 40, delete "an".

Col. 340, line 47, delete "an".

In the Sequence Listing

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Delete the Sequence Listing beginning in Col. 299, beginning with the text "<160> NUMBER OF SEQ ID NOS: 72" to and ending "<400> SEQUENCE: 72
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in Col. 340 and insert the following Sequence Listing:

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                                                                    96
Glu Asn Phe Lys Ala Leu Val Leu Ile Ala Phe Ala Gln Tyr Leu Gln
             20
                                  25
                                                      30
                                                                    144
caq tgt cca ttt gaa gat cat gta aaa tta gtg aat gaa gta act gaa
Gln Cys Pro Phe Glu Asp His Val Lys Leu Val Asn Glu Val Thr Glu
         35
ttt gca aaa aca tgt gtt gct gat gag tca gct gaa aat tgt gac aaa
                                                                    192
Phe Ala Lys Thr Cys Val Ala Asp Glu Ser Ala Glu Asn Cys Asp Lys
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	gaa Glu														288
	aga Arg														336
	cga Arg														384
	aat Asn 130														432
_	cat His				_	_	_					_			480
	aaa Lys	_	_		_	_	_		_	_	_		-	_	528
_	ctg Leu	_		_	_	_				_		_	-		576
	gcc Ala		_	_		_	_	_							624
	gct Ala 210														672
	gct Ala														720
	cac His														768
	gcg Ala														816

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											cct Pro	912
											tat Tyr	960
											gca Ala 335	1008
											aag Lys	1056
											cat His	1104
											gag Glu	1152
_				_					_		gga Gly	1200
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		Pro	Thr	Val	Glu	Val	Ser	Arg	Asn	Leu	gga Gly	1296
											ccc Pro	1344
											ttg Leu	1392
											gag Glu	1440

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						aag Lys		_			_				_	1584
						cac His 535										1632
	_	_	_	_	_	ttc Phe	-	_		_		_	_	_	_	1680
1 -	_	_	_			tgc Cys		_							_	1728
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Glu	Asn	Phe	Lys 20	Ala	Leu	Val	Leu	Ile 25	Ala	Phe	Ala	Gln	Tyr 30	Leu	Gln	
Gln	Cys	Pro 35	Phe	Glu	Asp	His	Val 40	Lys	Leu	Val	Asn	Glu 45	Val	Thr	Glu	
Phe	Ala 50	Lys	Thr	Cys	Val	Ala 55	Asp	Glu	Ser	Ala	Glu 60	Asn	Cys	Asp	Lys	
Ser 65	Leu	His	Thr	Leu	Phe 70	Gly	Asp	Lys	Leu	Cys 75	Thr	Val	Ala	Thr	Leu 80	
Arg	Glu	Thr	Tyr	Gly 85	Glu	Met	Ala	Asp	Cys 90	Cys	Ala	Lys	Gln	Glu 95	Pro	

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Glu	Arg	Asn	Glu 100	Cys	Phe	Leu	Gln	His 105	Lys	Asp	Asp	Asn	Pro 110	Asn	Leu
Pro	Arg	Leu 115	Val	Arg	Pro	Glu	Val 120	Asp	Val	Met	Cys	Thr 125	Ala	Phe	His
Asp	Asn 130	Glu	Glu	Thr	Phe	Leu 135	Lys	Lys	Tyr	Leu	Tyr 140	Glu	Ile	Ala	Arg
Arg 145	His	Pro	Tyr	Phe	Tyr 150	Ala	Pro	Glu	Leu	Leu 155	Phe	Phe	Ala	Lys	Arg 160
Tyr	Lys	Ala	Ala	Phe 165	Thr	Glu	Cys	Cys	Gln 170	Ala	Ala	Asp	Lys	Ala 175	Ala
Cys	Leu	Leu	Pro 180	Lys	Leu	Asp	Glu	Leu 185	Arg	Asp	Glu	Gly	Lys 190	Ala	Ser
Ser	Ala	Lys 195	Gln	Arg	Leu	Lys	Cys 200	Ala	Ser	Leu	Gln	Lys 205	Phe	Gly	Glu
Arg	Ala 210	Phe	Lys	Ala	Trp	Ala 215	Val	Ala	Arg	Leu	Ser 220	Gln	Arg	Phe	Pro
Lys 225	Ala	Glu	Phe	Ala	Glu 230	Val	Ser	Lys	Leu	Val 235	Thr	Asp	Leu	Thr	Lys 240
Val	His	Thr	Glu	Cys 245	Cys	His	Gly	Asp	Leu 250	Leu	Glu	Cys	Ala	Asp 255	Asp
Arg	Ala	Asp	Leu 260	Ala	Lys	Tyr	Ile	Cys 265	Glu	Asn	Gln	Asp	Ser 270	Ile	Ser
Ser	Lys	Leu 275	Lys	Glu	Cys	Cys	Glu 280	Lys	Pro	Leu	Leu	Glu 285	Lys	Ser	His
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				325		Leu			330					335	
Arg	His	Pro	Asp 340	Tyr	Ser	Val	Val	Leu 345	Leu	Leu	Arg	Leu	Ala 350	Lys	Thr
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Gln Val Ser Thr Pro Thr Leu Val Glu Val Ser Arg Asn Leu Gly Lys
            420
                                425
Val Gly Ser Lys Cys Cys Lys His Pro Glu Ala Lys Arg Met Pro Cys
                            440
Ala Glu Asp Tyr Leu Ser Val Val Leu Asn Gln Leu Cys Val Leu His
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Glu Lys Thr Pro Val Ser Asp Arg Val Thr Lys Cys Cys Thr Glu Ser
                                         475
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                                    490
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                                505
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Lys Ala Val Met Asp Asp Phe Ala Ala Phe Val Glu Lys Cys Cys Lys
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                  5
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tttagctcgg cttactcgag gggtgtgttt cgtcgagatg cacacaagag tgag
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Gly Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser
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aattcgaggg tgcaccgtca gtcttcctct tccccccaaa acccaaggac accctcatga
                                                                      120
                                                                      180
teteceggae teetgaggte acatgegtgg tggtggaegt aagecaegaa gaecetgagg
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tcaagttcaa ctggtacgtg gacggcgtgg aggt	gcataa tgccaagaca aagccgcggg 240						
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catcccggga tgagctgacc aagaaccagg tcag	cctgac ctgcctggtc aaaggcttct 480						
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site (GAS) containing promoter element
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aaatatctgc catctcaatt agtcagcaac catagtcccg cccctaactc cgcccatccc
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gcccctaact ccgcccaftt ccgcccattc tccgccccat ggctgactaa tttttttat
                                                                       180
ttatgcagag gccgaggccg cctcggcctc tgagctattc cagaagtagt gaggaggctt
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```
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<211> 73
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<213> Artificial Sequence
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<221> primer bind
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promoter element
<400> 81
gcggcctcga ggggactttc ccggggactt tccggggact ttccatcctg
                                                                     60
                                                                     73
ccatctcaat tag
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                                                                     60
caattagtca gcaaccatag tcccgccct aactccgccc atcccgccc taactccgcc
                                                                     120
cagttccgcc cattctccgc cccatggctg actaattttt tttatttatg cagaggccga
                                                                     180
ggccgcctcg gcctctgagc tattccagaa gtagtgagga ggcttttttg gaggcctagg
                                                                     240
cttttgcaaa aagctt
                                                                     256
```

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UNITED STATES PATENT AND TRADEMARK OFFICE CERTIFICATE OF CORRECTION

PATENT NO.

6,926,898

Page 1 of 35

APPLICATION NO.:

09/832,929

ISSUE DATE:

August 9, 2005

INVENTOR(S):

Craig A. Rosen and William A. Haseltine

It is hereby certified that an error or errors appear in the above-identified patent and that said Letters Patent is hereby corrected as shown below:

Under item (60) (Related U.S. Application Data) of the title page, delete the text beginning with "Provisional application No. 60/256,931" to and ending "provisional application No. 60/229,358, filed on Apr. 12, 2000."

Under item (56) (References Cited) of the title page and under U.S. PATENT DOCUMENTS beginning on page 1, insert:

	2003-0022308 A1	1/2003	Fleer et al.
÷	2003-0036170 A1	2/2003	Fleer et al.
	2003-0036171 A1	2/2003	Fleer et al.
	2003-0036172 A1	2/2003	Fleer et al.
	2003-0054554 A1	3/2003	Becquart et al.
	2003-0082747 A1	5/2003	Fleer et al.
	2003-0104578 A1	10/2001	Ballance
	2004-0010134 A1	4/2001	Rosen et al.
	09/832,501	4/2001	Ballance et al.
	09/833,041	4/2001	Rosen et al.
	09/833,111	4/2001	Rosen et al.
	09/833,117	4/2001	Rosen et al.
	09/833,118	4/2001	Rosen et al.
	10/702,536	11/2003	Fleer et al.
	10/702,636	11/2003	Fleer et al

MAILING ADDRESS OF SENDER

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Under item (56) (References Cited) of the title page and under OTHER PUBLICATIONS beginning on page 1, insert:

-- Larsson, M., et al., "Role of Annexins in Endocytosis of Antigens in Immature Human Dendritic Cells," *Immunology* 92:501-511 (1997).

Latta, M. et al., "Synthesis and Purification of Mature Human Serum Albumin From E. Coli," Bio/Technology 5:1309-1314 (1987).

Latta, M., et al., "Tryptophan Promoter Derivatives on Multicopy Plasmids: A Comparative Analysis of Expression Potentials in *Escherichia coli*," *DNA and Cell Biology* 9:129-137 (1990).

Lawn, R.M., et al., "The Sequence of Human Serum Albumin cDNA and its Expression in E. coli," *Nucleic Acids Research* 9:6103-6113 (1981).

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Leblois, H., et al., "Stable Transduction of Actively Dividing Cells via a Novel Adenoviral/Episomal Vector," *Molecular Therapy* 1:314-322 (2000).

Lee, C-H., et al., "Sodium Pertechnetate Tc99m Antral Scan in the Diagnosis of Retained Gastric Antrum," Arch. Surg. 119: 309-311 (1984).

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Lee, W-C., et al., "Identification and Characterization of a Nuclear Localization Sequence-Binding Protein in Yeast," *Proc. Natl. Acad. Sci. USA* 86:8808-8812 (1989).

Lee, Y-H., et al., "Comparison of Effective Renal Plasma Flow (ERPF) and Endogenous Creatinine Clearance (Ccr) in Evaluation of the Differential Kidney Function: An in Vivo Study," *Chin. Med. J. (Taipei)* 49:147-152 (1992).

Lei, H-Y., et al., "An Antigen-specific Hypersensitivity Which Does Not Fit Into Traditional Classification of Hypersensitivity," *The Journal of Immunology* 143:432-438 (1989).

Levitt, D., et al., "Toxicity of Perfluorinated Fatty-Acids for Human and Murine B Cell Lines," *Toxicology and Applied Pharmacology* 86:1-11 (1986).

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Lew D.B., et al., "Mitogenic Effect of Lysosomal Hydrolases on Bovine Tracheal Myocytes in Culture," *The Journal of Clinical Investigation* 88:1969-1975 (1991).

Lewis, C., et al., "Is Sexual Dysfunctoin in Hypertensive Women Uncommon or Understudied?" American Jour of Hypertension," 11:733-735 (1998). --

Under item (57) (ABSTRACT) of the title page, "disordrs" should read --disorders--.

In the Specification

Col. 143, line 26, delete "As exhibited in Table 2, most", and insert -- Most--.

Col. 143, line 31, delete "Table 2".

In the Claims

Col. 340, line 40, delete "an".

Col. 340, line 47, delete "an".

In the Sequence Listing

Delete the Sequence Listing beginning in Col. 299, beginning with the text "<160> NUMBER OF SEQ ID NOS: 72" to and ending "<400> SEQUENCE: 72

Gly Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser

in Col. 340 and insert the following Sequence Listing:

<160> NUMBER OF SEQ ID NOS: 82

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<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<221> primer bind

<223> primer useful to clone human growth hormone cDNA

<400> 1

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23

<210> 2

<211> 33

<212> DNA

<213> Artificial Sequence

<220>

<221> primer_bind

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<220>
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<223> synthetic oligonucleotide used to join DNA fragments with non-cohesive
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Ile Ser Ala Asp Ala His Lys Ser
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Finnegan, Henderson, Farabow, Garrett & Dunner, L.L.P. 901 New York Avenue, N.W. Washington, D.C. 20001-4413

JAN 04 2006

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<211> 47
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Asp Ala His Lys Ser Glu Val Ala His Arg Phe Lys Asp Leu Gly Glu
gaa aat ttc aaa gcc ttg gtg ttg att gcc ttt gct cag tat ctt cag
                                                                    96
Glu Asn Phe Lys Ala Leu Val Leu Ile Ala Phe Ala Gln Tyr Leu Gln
cag tgt cca ttt gaa gat cat gta aaa tta gtg aat gaa gta act gaa
                                                                    144
Gln Cys Pro Phe Glu Asp His Val Lys Leu Val Asn Glu Val Thr Glu
                             40.
ttt gca aaa aca tgt gtt gct gat gag tca gct gaa aat tgt gac aaa
                                                                   192
Phe Ala Lys Thr Cys Val Ala Asp Glu Ser Ala Glu Asn Cys Asp Lys
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Arg Glu Thr Tyr Gly Glu Met Ala Asp Cys Cys Ala Lys Gln Glu Pro 85 gag aga aat gaa tgc ttc ttg caa cac aaa gat gac aac cca aac ctc Glu Arg Asn Glu Cys Phe Leu Gln His Lys Asp Asp Asn Pro Asn Leu 100 ccc cga ttg gtg aga cca gag gtt gat gtg atg tgc act gct ttt cat 110 ccc cga ttg gtg aga cca gag gtt gat gtg atg tgc act gct ttt cat 115 115 120 gac aat gaa gag aca ttt ttg aaa aaa tac tta tat gaa att gcc aga Asp Asn Asn Glu Glu Thr Phe Leu Lys Lys Tyr Leu Tyr Glu Ile Ala Arg 130 aga cat gat gat gac ccc ggaa ctc ctt ttc ttt gct aaa agg 480 Arg His Pro Tyr Phe Tyr Ala Pro Glu Leu Leu Phe Phe Ala Lys Arg 145 150 tat aaa gct gct ttt aca gaa tgt tgc caa gct gct gat aaa gct gcc Tyr Lys Ala Ala Phe Thr Glu Cys Cys Gln Ala Ala Asp Lys Ala Ala 165 tgc ctg ttg cca aag ctc gat gaa ctt ctg ggat gaa ggg ag gct tcg Cys Leu Leu Pro Lys Leu Asp Glu Leu Arg Asp Glu Cly Lys Ala Ser 180 tct gcc aaa cag aga ctc aaa tgt gcc agt ctc caa aaa ttt gga gaa Ser Ala Lys Gln Arg Leu Lys Cys Ala Ser Leu Gln Lys Phe Gly Glu 195 aga gct ttc aaa gca tgg gca gtg gct cgc ctg agc cag aga ttt ccc Arg Ala Phe Lys Ala Trp Ala Val Ala Arg Leu Ser Gln Arg Phe Pro 210 210 aaa gct gag ttt gca gaa gtt cc aag tta gtg aga gat gct gct gat aaa gct gcc G72 Arg Ala Glu Phe Ala Glu Val Ser Lys Leu Val Thr Asp Leu Thr Lys 225 agg gcg gac ctt gcc aag tat atc tgt gaa aat cag gat tcg gat gac G768 Val His Thr Glu Cys Cys His Gly Asp Leu Leu Glu Cys Ala Asp
Glu Arg Asn Glu Cys Phe Leu Gln His Lys Asp Asp Asn Pro Asn Leu 100 110 110 ccc cga ttg gtg aga cca gag gtt gat gtg atg tgc act gct ttt cat 384 Pro Arg Leu Val Arg Pro Glu Val Asp Val Met Cys Thr Ala Phe His 120 125 gac aat gaa gag aca ttt ttg aaa aaa tac tta tat gaa att gcc aga 432 Asp Asn Glu Glu Thr Phe Leu Lys Lys Tyr Leu Tyr Glu Ile Ala Arg 130 135 140 aga cat cct tac ttt tat gcc ccg gaa ctc ctt ttc ttt gct aaa agg 480 Arg His Pro Tyr Phe Tyr Ala Pro Glu Leu Leu Phe Phe Ala Lys Arg 160 tat aaa gct gct ttt aca gaa tgt tgc caa gct gct gat aaa gct gcc Tyr Lys Ala Ala Phe Thr Glu Cys Cys Gln Ala Ala Asp Lys Ala Ala 165 170 tgc ctg ttg cca aag ctc gat gaa ctt cgg gat gaa ggg aag gct tcg Cys Leu Leu Pro Lys Leu Asp Glu Leu Arg Asp Glu Gly Lys Ala Ser 180 tct gcc aaa cag aga ctc aaa tgt gcc agt ctc caa aaa ttt gga gaa gct tcg Cys Leu Leu Pro Lys Leu Asp Glu Leu Arg Asp Glu Gly Lys Ala Ser 180 tct gcc aaa cag aga ctc aaa tgt gcc agt ctc caa aaa ttt gga gaa Get Ser Ala Lys Gln Arg Leu Lys Cys Ala Ser Leu Gln Lys Phe Gly Glu 205 aga gct ttc aaa gca tgg gca gtg gct cgc ctg agc cag aga ttt ccc Arg Ala Phe Lys Ala Trp Ala Val Ala Arg Leu Ser Gln Arg Phe Pro 210 225 aaa gct gag ttt gca gaa gtt tcc aag tta gtg aca gat ctt acc aaa 125 Cys Ala Glu Phe Ala Glu Val Ser Lys Leu Val Thr Asp Leu Thr Lys 225 230 240 gcc cac acg gaa tgc tgc cat gga gat ctg ctt gaa tgt gct gat gac Val His Thr Glu Cys Cys His Gly Asp Leu Leu Glu Cys Ala Asp Asp 245 agg gcg gac ctt gcc aag tat atc tgt gaa aat cag gat tcg atc cc Arg Ala Aps Leu Ala Aps Asp Leu Ala Ala Aps Cag Glu Asp Cu Leu Cu Glu Cys Ala Asp Asp 245 agg gcg gac ctt gcc aag tat atc tgt gaa aat cag gat tcg atc tcc Arg Ala Aps Leu Ala Ala Aps Asp Cau Ala Ala Aps Asp Leu Ala Ala Aps Asp Cau Ala Ala Ala Che Arg Ala Aps Asp Cau Ala Ala Ala Che Arg Ala Aps Leu Ala Che Arg Ala Aps Asp Cau Ala Ala Ala Che Arg Ala Aps Asp Cau Ala Ala Ala Che A
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Asp Asn Glu Glu Thr Phe Leu Lys Lys Tyr Leu Tyr Glu Ile Ala Arg 130 135 140 140 140 140 140 135 140 140 135 140 140 135 140 140 140 140 140 140 140 140 140 140
Arg His Pro Tyr Phe Tyr Ala Pro Glu Leu Leu Phe Phe Ala Lys Arg 150
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Val His Thr Glu Cys Cys His Gly Asp Leu Leu Glu Cys Ala Asp Asp 245 250 255 agg gcg gac ctt gcc aag tat atc tgt gaa aat cag gat tcg atc tcc Arg Ala Asp Leu Ala Lys Tyr Ile Cys Glu Asn Gln Asp Ser Ile Ser
Arg Ala Asp Leu Ala Lys Tyr Ile Cys Glu Asn Gln Asp Ser Ile Ser

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			_														
_	aaa Lys	Leu	-		_	_	Glu			_		Glu				864	<i>:</i> .
_	att.	_	_		_		_		-		_	-	_			912	
Cys	Ile 290	Ala	Glu	Val	Glu	Asn 295	Asp	Glu	Met	Pro	Ala 300	Asp	Leu	Pro	Ser		
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	aaa Lys					Āsp										1440	

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Glu	Asn	Phe	Lys 20	Ala	Leu	Val	Leu	Ile 25	Ala	Phe	Ala	Gln	Tyr 30	Leu	Gln			
Gln	Cys	Pro 35	Phe	Glu	Asp	His	Val 40	Lys	Leu	Val	Asn	Glu 45	Val	Thr	Glu			
Phe	Ala 50	Lys	Thr	Cys	Val	Ala 55	Asp	Glu	Ser	Ala	Glu 60	Asn	Cys	Asp	Lys			
Ser 65	Leu	His	Thr	Leu	Phe 70	Gly	Asp	Lys	Leu	Cys 75	Thr	Val	Ala	Thr	Leu 80			
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	•													_	
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Asp	Asn 130	Glu	Glu	Thr	Phe	Leu 135	Lys	Lys	Tyr	Leu	Tyr 140	Glu	Ile	Ala	Arg
Arg 145	His	Pro	Tyr	Phe	Tyr 150	Ala	Pro	Glu	Leu	Leu 155	Phe	Phe	Ala	Lys	Arg 160
Tyr	Lys	Ala	Ala	Phe 165	Thr	Glu	Cys	Cys	Gln 170	Ala	Ala	Asp	Lys	Ala 175	Ala
Cys	Leu	Leu	Pro 180	Lys	Leu	Asp	Glu	Leu 185	Arg	Asp	Glu		Lys 190	Ala	Ser
Ser	Ala	Lys 195	Gln	Arg	Leu	Lys	Cys 200	Ala	Ser	Leu	Gln	Lys 205	Phe	Gly	Glu
Arg	Ala 210	Phe	Lys	Ala	Trp	Ala 215	Val	Ala	Arg		Ser 220	Gln	Arg	Phe	Pro
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Val	His	Thr	Glu	Cys 245	Cys	His	Gly	Asp	Leu 250	Leu	Glu	Cys	Ala	Asp 255	Asp
Arg	Ala	Asp	Leu 260	Ala	Lys	Tyr		Cys 265	Glu	Asn	Gln	Asp	Ser 270	Ile	Ser
Ser	Lys	Leu 275	Lys	Glu	Cys	Cys	Glu 280	Lys	Pro	Leu	Leu	Glu 285	Lys	Ser	His
Cys	Ile 290	Ala	Glu	Val	Glu	Asn 295	Asp	Glu	Met	Pro	Ala 300	Asp	Leu	Pro	Ser
Leu 305	Ala	Ala	Asp	Phe	Val 310	Glu	Ser	Lys	Asp	Val 315	Cys	Lys	Asn	Tyr	Ala 320
Glu	Ala	Lys	Asp	Val 325		Leu	Gly	Met	Phe 330	Leu	Tyr	Glu	Tyr	Ala 335	Arg
Arg	His	Pro	Asp 340	Tyr	Ser	Val	Val	Leu 345	Leu	Leu	Arg	Leu	Ala 350	Lys	Thr
Tyr	Glu	Thr 355	Thr	Leu	Glu	Lys	Cys 360	Cys	Ala	Ala	Ala	Asp 365	Pro	His	Glu
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Gln Asn Leu Ile Lys Gln Asn Cys Glu Leu Phe Glu Gln Leu Gly Glu
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Tyr Lys Phe Gln Asn Ala Leu Leu Val Arg Tyr Thr Lys Lys Val Pro
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Gln Val Ser Thr Pro Thr Leu Val Glu Val Ser Arg Asn Leu Gly Lys
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Val Gly Ser Lys Cys Cys Lys His Pro Glu Ala Lys Arg Met Pro Cys
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Leu Val Asn Arq Arq Pro Cys Phe Ser Ala Leu Glu Val Asp Glu Thr
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Ile Cys Thr Leu Ser Glu Lys Glu Arg Gln Ile Lys Lys Gln Thr Ala
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Leu Val Glu Leu Val Lys His Lys Pro Lys Ala Thr Lys Glu Gln Leu
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tcaagttcaa ctggtacgtg gacggcgtgg aggtgcataa tgccaagaca aagccgcg	gg 240
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ggctgaatgg caaggagtac aagtgcaagg tctccaacaa agccctccca acccccat	cg 360
agaaaaccat ctccaaagcc aaagggcagc cccgagaacc acaggtgtac accctgcc	cc 420
catcccggga tgagctgacc aagaaccagg tcagcctgac ctgcctggtc aaaggctt	ct 480
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acaagagcag gtggcagcag gggaacgtct tetcatgete egtgatgcat gaggetet	gc 660
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gcccctaact ccgcccagtt ccgcccattc tccgccccat ggctgactaa ttttttttat
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ttatgcagag gccgaggccg cctcggcctc tgagctattc cagaagtagt gaggaggctt
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